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	CLASS	SUBCLASS

Notch(C)	IDE <u>C</u> -SNP	C <u>Q</u> NGGTC---D-V <u>Q</u> GSY- <u>C</u> -CPP <u>Q</u> GFT	G <u>Q</u> K---CE-N
<u>10244(C)</u>	-NE <u>Q</u> CTM---	<u>C</u> QH--- <u>C</u> VNT- <u>Q</u> SY- <u>C</u> KG- <u>Q</u> SG- <u>C</u> CRCFP <u>Q</u> GYT	G---I- <u>C</u> D
42		<u>C</u> GKT	<u>C</u> SQ D
57	VNE <u>Q</u> CGMKPRP	<u>C</u> QHR	<u>C</u> GHLMP D
108	VNSRTCAMIN	<u>C</u> QYS	<u>C</u> GLRLAPN
166	IDE <u>Q</u> CASGKVI	<u>C</u> PYNRR <u>C</u>	<u>C</u> QYISGR
212	IN <u>Q</u> ECTMD <u>Q</u> HT	<u>C</u> SHH <u>Q</u> ANG	<u>C</u> GNGRLCS
CD97(C)	V- <u>Q</u> E <u>Q</u> -SG- <u>Q</u> -- <u>C</u> -SS- <u>C</u>	-NTV <u>Q</u> GSY- <u>C</u> R <u>Q</u> RP <u>Q</u> W-P-P <u>Q</u> -PN---	D
EGF(C)	NSDSEC <u>Q</u> PLSHDG <u>Q</u> YCLHDGV <u>Q</u> MYIEALDKYACNCVVG <u>Q</u> YI---GER--- <u>C</u> QYRDLKWWELR		

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	CLASS SUBCLASS

GGCTGGAGAA GAAACAGCAA GGGAGTCTGT GAAGCTACAT GCGAACCTGG
 ATGTAAGTTT GGTGAGTGC G TGGACCAAA CAAATGCAGA TGCTTCAG
 GATACACCGG GAAAACCTGC AGTCAAGATG TGAATGAGTG TGGATGAAA
 CCCCCGCCAT GCCAACACAG ATGTGTGAAT ACACACGGAA GCTACAAGTG
 CTTTGCCTC AGTGGCCACA TGCTCATGCC AGATGCTACG TGTGTGAAC
 CNAGGACATG TGCCATGATA AACTGTCAGT ATAGCTGTGA AGACACAGAA
(SEQ ID NO 1)

GGCTGGAGAA GAAACAGCAA GGGAGTCTGT GAAGCTACAT GCGAACCTGG
 ATGTAAGTTT GGTGAGTGC G TGGACCAAA CAAATGCAGA TGCTTCAG
 GATACACCGG GAAAACCTGC AGTCAAGATG TGAATGAGTG TGGATGAAA
 CCCCCGCCAT GCCAACACAG ATGTGTGAAT ACACACGGAA GCTACAAGTG
 CTTTGCCTC AGTGGCCACA TGCTCATGCC AGATGCTACG TGTGTGAAC
 CNAGGACATG TGCCATGATA AACTGTCAGT ATAGCTGTGA AGACACAGAA
 GAAGGGCCAC AGTGCCTGTG TCCATCCTCA GGACTCCGCC TGGCCCCAAA
 TGAAGAGAC TGTCTAGATA TTGATGAATG TGCCTCTGGT AAAGTCATCT
 GTCCCTACAA TCGAAGATGT GTGAACACAT TTGGAAGCTA CTACTGCAAA
 TGTACATTG GTTCTGAAC GCAATATATC AGTGGACGAT ATGACTGTAT
 AGATATAAT GAATGTACTA TGGATAGCCA TACGTGCAGC CACCATGCCA
 ATTGCTTCAA TACCCAAGGG TCCTTCAAGT GTAAATGCAA GCAGGGATAT
 AAAGGCAATG GACTTCGGTG TTCTGCTATC CCTGAAAATT CTGTGAAGGA
 AGTCCTCAGA GCACCTGGTA CCATCAAAGA CAGAATCAAG AAGTTGCTTG
 CTCACAAAAA CAGCATGAAA AAGAAGGCAA AAATAAAAAA TGTTACCCCA
 GAACCCACCA GGACTCCTAC CCCTAAGGTG AACTTGCAGC CCTTCAACTA
 TGAAGAGATA GTTCCAGAG GCGGGAACTC TCATGGAGGT AAAAAAGGGA
 ATGAAGAGAA AATGAAAGAG GGGCTTGAGG ATGAGAAAAG AGAAGAGAAA
 GCCCTGAAGA ATGACATAGA GGAGCGAAGC CTGCGAGGAG ATGTGTTTT
 CCCTAAGGTG AATGAAGCAG GTGAATTGG C CTGATTCTG GTCCAAAGGA
 AAGCGCTAAC TTCCAAACTG GAACATAAAG ATTTAAATAT CTCGGTTGAC
 TGCAGCTTCA ATCATGGAT CTGTGACTGG AAACAGGATA GAGAAGATGA
 TTTGACTGG AATCCTGCTG ATCGAGATAA TGCTATTGGC TTCTATATGG
 CAGTTCCGGC CTTGGCAGGT CACATGAAAG ACATTGGCCG ATTGAAACTT
 CTCTACCTG ACCTGCAACC CCAAAGCAAC TTCTGTTGC TCTTGATTA
 CCGGCTGGCC GGAGACAAAG TCGGGAAACT TCGAGTGTG GTAAAAAGCA
 GTAACAATGC CCTGGCATGG GAGAAGACCA CGAGTGAGGA TGAAAAGTGG
 AAGACAGGGA AAATTCAGTT GTATCAAGGA ACTGATGCTA CCAAAAGCAT
 CATTGGAA GCAGAACGTG GCAAGGGCAA AACCGGCGAA ATCGCAGTGG
 ATGGCGTCTT GCTTGTTC G GCTTATGTC CAGATAGCCT TTTATCTGTG
 GANNNCTGAA TGGTACTATC TTTATATTG ACTTTGTATG TCAGTTCCCT
 GGTTTTTG ATATTGCATC ATAGGACCTC TGGCATTAA AAATTACTAG
 CTGAAAAATT G
(SEQ ID NO 2)

FIG. 2

APPROVED BY DRAFTSMAN	O.G. FIG. CLASS	SUBCLASS
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GWRRNSKGVEATCEPGCKFGEVGPNKCRCFPGYTGKTCSDVNECGMKPRPCQHR
 CVNTHGSYKCFCLSGHMLMPDATCVNSRTCAMINCQYSCEDTEEGPQCLCPSSGLRLAP
(SEQ ID NO 3)

GWRRNSKGVEATCEPGCKFGEVGPNKCRCFPGYTGKTCSDVNECGMKPRPCQHR
 CVNTHGSYKCFCLSGHMLMPDATCVNSRTCAMINCQYSCEDTEEGPQCLCPSSGLRLAP
 NGRDCLDIDEASGKVICPYNRRCVNTFGSYYCKCHIGFELQYISGRYDCIDINECTMDS
 HTCSHHANCNTQGSFKCKCKQGYKGNGLRCSAIPENSKEVLRAPGTIKDRIKKLLAH
 KNSMKKAKIKNVTPEPTRTPKVNLPNQPFNYEEIVSRGGNSHGGKKGNEEKMKEGLE
 DEKREEKALKNDIEERSLRGDVFFPKVNEAGEFGLILVQRKALTSLKLEHKDLNISVDCSF
 NHGICDWKQDREDDFDWNPADRNAIGFYMAVPALAGHMKDIGRLKLLPDLQPQSN
 FCLLFDYRLAGDKVGKLRVFKNSNNALAWEKTTSEDEKWKTGKIQLYQGTDATKSIIF
 EAERGKGKTGEIAVDGVLLVSGLCPDSLLSVDDXMVLSLYLTLVSSLVFLILHHRTSGI
 LKLLAEKL
(SEQ ID NO 4)

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FIG. 3